

(FILE 'HOME' ENTERED AT 11:37:05 ON 12 FEB 2004)

FILE 'MEDLINE, EMBASE, BIOSIS' ENTERED AT 11:37:25 ON 12 FEB 2004

L1 0 S "NEED ACTIVATION DOMAIN" (P) CHROMATIN
L2 254 S "ACTIVATION DOMAIN" (P) CHROMATIN
L3 85 S L2 (P) REMODELING
L4 31 DUP REM L3 (54 DUPLICATES REMOVED)
L5 3 S L2 AND "GENE SPECIFIC"
L6 3168 S "CHROMATIN REMODELING"
L7 13 S L6 (S) "GENE SPECIFIC"
L8 13 S L7 (P) "GENE SPECIFIC"
L9 28 S L6 (P) "GENE SPECIFIC"
L10 28 S L6 AND "GENE SPECIFIC"
L11 9 DUP REM L7 (4 DUPLICATES REMOVED)
L12 11 DUP REM L9 (17 DUPLICATES REMOVED)
L13 11 DUP REM L10 (17 DUPLICATES REMOVED)
L14 3974 S SWI/SNF OR RSC OR NURD OR NURF OR ACF OR CHRAC OR RSF OR ISW1
L15 5108 S "SWI/SNF" OR RSC OR NURD OR NURF OR ACF OR CHRAC OR RSF OR IS
L16 18 S L15 (S) "GENE SPECIFIC"
L17 9 DUP REM L16 (9 DUPLICATES REMOVED)
L18 18910 S "ZINC FINGER"
L19 0 S L18 (S) "CHROMATIN REMODEL"
L20 0 S L18 (P) "CHROMATIN REMODEL"
L21 0 S L18 AND "CHROMATIN REMODEL"
L22 545 S L18 AND CHROMATIN
L23 0 S L18 (P) CHROMTAIN
L24 440 S L18 (P) CHROMATIN
L25 191 S L18 (S) CHROMATIN
L26 135 DUP REM L25 (56 DUPLICATES REMOVED)
L27 66 S L26 NOT PY>=2001
L28 178 DUP REM L24 (262 DUPLICATES REMOVED)
L29 1 DUP REM L5 (2 DUPLICATES REMOVED)
L30 135 DUP REM L25 (56 DUPLICATES REMOVED)
L31 80 S L28 NOT PY>=2001
L32 1 S L29 NOT PY>=2001
L33 15 S L31 AND FUSION
L34 8 S TARGETED (P) ("CHROMATIN ALTERATION" OR "ALTERED CHROMATIN" O
L35 4 DUP REM L34 (4 DUPLICATES REMOVED)